Changed the margins in cases where the sequence text was "wapped" downto the lext line.  Edited a lormatierror in the Current Application Data section, specifically:  Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other applicant spelled out a number instead of using an integer changed the "Number of Sequences" field. The applicant spelled out a number instead of using an integer changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEO ID NO when obviously incorrect. The sequence gumbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: [In non-ASCII 'garbage' at the beginning fend of files; secretary initials/filename at end or page numbers throughout text; other invalid text, such as inserted an obvious error in the response, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted.		chor: 09/781 1/8 ENTERE Strick of the day: 12/1/08/2009 (8TIC)
Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other  Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer Changed the spelling of a mandatory field (the headings of subheadings), specificands.  Corrected the SEO ID NO when obviously incorrect. The sequence gumbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted:	Cha	inged the margins in cases where the sequence text was "wrapped" down-to the next line.
Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integround of the spelling of a mandatory field (the headings of subheadings), specificary:  Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Defeted extra, invalid, headings used by an applicant, specifically:  Defeted extra, invalid, headings used by an applicant, specifically:  Corrected an obvious error in the response, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an orror in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be defeted.	Edit	ed a format error in the Current Application Data section, specifically:
Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integral Changed the spelling of a mandatory field (the headings of subheadings), specifically:  Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Defeted extra, invalid, headings used by an applicant, specifically:  Defeted: Innon-ASCII 'garbage' at the beginning/end of files: secretary initials/filename at end of page numbers throughout text; other invalid text, such as  Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be defeted.	 Edit app	ed the Current Application Data section with the actual current number. The number inputted by the licant was the prior application data; or other
Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: One-ASCII garbage at the beginning and of files: secretary initials/filename at end one page numbers throughout text; other invalid text, such as  Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted.	 Add	ed the mandatory heading and subheadings for "Current Application Data".
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Inserted colons after headings/subheadings. Headings edited included:  Defeted extra, invalid, headings used by an applicant, specifically:  Defeted: non-ASCII "garbage" at the beginning/end of files: secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be defeted.	Inse	rtod or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:
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Deleted: In non-ASCII "garbage" at the beginning/end of files: secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Insc	erted colons after headings/subheadings. Headings edited included:
Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Dole	oted extra, invalid, headings used by an applicant, specifically:
Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be defeted.  Defeted and final stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (en	Del	etod: non-ASCII "garbago" at the beginning/end of files; secretary initials/filename at end of files page numbers throughout text; other invalid text, such as
Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be defeted.  Defeted and logistical stop coden in amino acid sequences and adjusted the "(A)Length:" field accordingly (en	Ins	orted mandatory headings, specifically:
Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be defeted.  Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (en	Coï	rected an obvious error in the response, specifically:
Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be defeted.  Deleted ending, stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (en	Edi	led identifiers where upper case is used but lower case is required, or vice versa.
A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (en		rected an error in the Number of Sequences field, specifically:
Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (enduced to a Patentin bug). Sequences corrected:	۱. ۷	•
	Delai	ed ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (erro
Other:		

LExaminer: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

RAW SEQUENCE LISTING DATE: 03/29/2001 PATENT APPLICATION: US/09/786,648 TIME: 11:00:23

Input Set : A:\Pto.da

7 <110> APPLICANT: Williams, Neil Andrew

Output Set: N:\CRF3\03292001\1786648.raw

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Hirst, Timothy Raymond
     11 <120> TITLE OF INVENTION: Peptide Fragments of Cholera Toxin B or Enterotoxin B as Vaccine
     12
              Adjuvants
     14 <130> FILE REFERENCE: 7438
C--> 16 <140> CURRENT APPLICATION NUMBER: US/09/786,648
     18 <141> CURRENT FILING DATE: 2001-03-07
     20 <150> PRIOR APPLICATION NUMBER: PCT/GB99/02970
     22 <151> PRIOR FILING DATE: 1999-09-07
     24 <160> NUMBER OF SEQ ID NOS: 6
     26 <170> SOFTWARE: MS DOS
     30 <210> SEQ ID NO: 1
     32 <211> LENGTH: 27
     34 <212> TYPE: DNA
     36 <213> ORGANISM: artificial sequence
     38 <220> FEATURE:
     40 <223> OTHER INFORMATION: encodes heptapeptide of SEQ ID NO 2
     42 <400> SEQUENCE: 1
     44 gaagtaccag gtagtcaaca tatagat
                                                           27
     48 <210> SEQ ID NO: 2
     50 <211> LENGTH: 7
     52 <212> TYPE: PRT
     54 <213> ORGANISM: E. coli
     56 <220> FEATURE:
    58 <222> LOCATION: 51...57
    60 <223> OTHER INFORMATION: isolated or synthetic EtxB beta4-alpha2 loop fragment derivable from
              human variant E. coli
    63 <400> SEQUENCE: 2
    65 Glu Val Pro Gly Ser Gln His
    70 <210> SEQ ID NO: 3
    72 <211> LENGTH: 12
    74 <212> TYPE: PRT
    76 <213> ORGANISM: E. coli
    78 <220> FEATURE:
    80 <222> LOCATION: 50...61
    82 <223> OTHER INFORMATION: isolated or synthetic EtxB beta4-alpha2 loop fragment derivable from
             human variant E. coli
    85 <400> SEQUENCE: 3
    87 Val Glu Val Pro Gly Ser Gln His Ile Asp Ser Gln
    92 <210> SEQ ID NO: 4
    94 <211> LENGTH: 21
    96 <212> TYPE: PRT
    98 <213> ORGANISM: E. coli
    100 <220> FEATURE:
    102 <222> LOCATION: 45...65
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TIME: 11:00:23

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/786,648

Input Set : A:\Pto.da

Output Set: N:\CRF3\03292001\I786648.raw

```
104 <223> OTHER INFORMATION: isolated or synthetic EtxB beta4-alpha2 loop fragment derivable from
        human variant E. coli
107 <400> SEQUENCE: 4
110 Gly Ala Thr Phe Gln Val Glu Val Pro Gly Ser Gln His Ile Asp
                      5
                                        10
113 Ser Gln Lys Lys Ala Ile
114
                     20
118 <210> SEQ ID NO: 5
120 <211> LENGTH: 21
122 <212> TYPE: PRT
124 <213> ORGANISM: E. coli
126 <220> FEATURE:
128 <222> LOCATION: 45...65
130 <223> OTHER INFORMATION: isolated or synthetic EtxB beta4-alpha2 loop fragment derivable from
         porcine E. coli
133 <400> SEQUENCE: 5
135 Gly Glu Thr Phe Gln Val Glu Val Pro Gly Ser Gln His Ile Asp
                    5
138 Ser Gln Lys Lys Ala Ile
139
143 <210> SEQ ID NO: 6
145 <211> LENGTH: 13
147 <212> TYPE: PRT
149 <213> ORGANISM: artificial sequence
151 <220> FEATURE:
153 <223> OTHER INFORMATION: control peptide used in Example 5
155 <400> SEQUENCE: 6
157 Leu Arg Asn Glu Thr Thr Thr Thr Lys Gly Asp Tyr Cys
                      5
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/786,648



DATE: 03/29/2001 TIME: 11:00:24

Input Set : A:\Pto.da

Output Set: N:\CRF3\03292001\I786648.raw

L:16 M:270 C: Current Application Number differs, Replaced Application Number

PCT

RAW SEQUENCE LISTING

DATE: 03/23/2001

PATENT APPLICATION: US/09/786,648

TIME: 09:44:37

Input Set : A:\es.txt

Output Set: N:\CRF3\03232001\1786648.raw

Does Not Comply
Corrected Diskette Needed

## ERRORED SEQUENCES

143 <210> SEQ ID NO: 6
145 <211> LENGTH: 13
147 <212> TYPE: PRT
149 <213> ORGANISM: artificial sequence
151 <220> FEATURE:
153 <223> OTHER INFORMATION: control peptide used in Example 5
155 <400> SEQUENCE: 6
157 Leu Arg Asn Glu Thr Thr Thr Lys Gly Asp Tyr Cys
158
5 10
E--> 159
??



VERIFICATION SUMMARY

PATENT APPLICATION: US/09/786,648

DATE: 03/23/2001 TIME: 09:44:38

Input Set :  $A:\es.txt$ 

Output Set: N:\CRF3\03232001\I786648.raw

 ${\tt L:16~M:270~C:} \ {\tt Current~Application~Number~differs,~Replaced~Application~Number}$ 

L:159 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:159 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1